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- (54) **ANALOGUES DE L'HORMONE PARATHYROIDIENNE
UTILISES POUR LE TRAITEMENT DE L'OSTEOPOROSE**
- (54) **PARATHYROID HORMONE ANALOGUES FOR THE
TREATMENT OF OSTEOPOROSIS**

(57) Certain analogues of human parathyroid hormone (hPTH) have been found to be effective for the treatment of osteoporosis, while showing decreased side effects. Analogues showing this effect include all sequences from hPTH-(1-28)-NH₂ to hPTH-(1-31)-NH₂ and all sequences from [Leu²⁷]-hPTH-(1-28)-NH₂ to [Leu²⁷]-hPTH-(1-34)-NH₂. Also included are cyclic analogues cyclo(Lys²⁶-Asp³⁰)[Leu²⁷]-hPTH-(1-34)NH₂ and cyclo(Lys²⁷-Asp³⁰)-hPTH-(1-34)-NH₂. Analogues in the form of the carboxyl terminal amide are particularly effective.

Abstract

Certain analogues of human parathyroid hormone (hPTH) have been found to be effective for the treatment of osteoporosis, while showing decreased side effects. Analogues showing this effect include all sequences from hPTH-(1-28)-NH₂ to hPTH-(1-31)-NH₂ and all sequences from [Leu²⁷]-hPTH-(1-28)-NH₂ to [Leu²⁷]-hPTH-(1-34)-NH₂. Also included are cyclic analogues cyclo(Lys²⁶-Asp³⁰)[Leu²⁷]-hPTH-(1-34)NH₂ and cyclo (Lys²⁷-Asp³⁰)-hPTH-(1-34)-NH₂. Analogues in the form of the carboxyl terminal amide are particularly effective.

Parathyroid Hormone Analogues for the Treatment of
Osteoporosis

Field of the Invention

This invention relates to analogues of human parathyroid
5 hormone, which have been found to be effective in the
treatment of osteoporosis and will reverse the loss of bone
and increase bone mass and density specifically without
undesirable side effects.

Background of the Invention

10 Osteoporosis is a leading cause of disability in the
elderly, particularly elderly women. It is well known that
human parathyroid hormone (hPTH) and certain analogues are
useful in the treatment of osteoporosis.

Parathyroid hormone (PTH) is produced by the parathyroid
15 gland and is involved in the control of calcium levels in
blood. It is a hypercalcemic hormone, elevating blood calcium
levels. PTH is a polypeptide and synthetic polypeptides
containing the first thirty-four residues of PTH may be
prepared using the method disclosed by Erickson and
20 Merrifield, The Proteins, Neurath et al., Eds., Academic
Press, New York, 1976, page 257, preferably as modified by the
method of Hodges et al., Peptide Research, 1, 19 (1988).

When serum calcium is reduced to below a "normal" level,
the parathyroid gland releases PTH and resorption of bone
25 calcium and increased absorption of calcium from the
intestine, as well as renal reabsorption of calcium, occur.
The antagonist of PTH is calcitonin, which acts to reduce the
level of circulating calcium. Osteoporosis is a progressive
disease which results in the reduction of total bone mass.
30 This often results in fractures of load-bearing bones and the
physical degenerations characteristic of immobilizing
injuries. Osteoporosis is associated with hyperthyroidism,
hyperparathyroidism, Cushing's syndrome and the use of certain
steroidal drugs. Remedies historically have involved increase
35 in dietary calcium, estrogen therapy and increased doses of
vitamin D.

Although high levels of PTH can remove calcium from the
bone, low doses can actually promote bone growth.

While the use of PTH is effective in the treatment of osteoporosis by diminishing the loss of bone mass, PTH may exhibit other undesired pharmacological effects, such as hypertension and smooth muscle relaxation (e.g. relaxation of gastrointestinal organs, uterus, tracheal and vas deferens) as well as positive chronotropic and inotropic effects on the heart.

Tregear U.S. Patent 4,086,196, describes human PTH analogues and claims that the first 27 to 34 amino acids are the most effective in terms of activation of adenylyl cyclase. Rosenblatt et al, U.S. Patent 4,771,124 discloses the property of hPTH analogues wherein Trp²³ is substituted by amino acids phenylalanine, leucine, norleucine, valine, tyrosine, beta-naphthylalanine and alpha-naphthylalanine as a PTH antagonist. These modified hPTH analogues also have the 2 and 6 amino terminal amino acids removed, resulting in loss of most agonist activities when used to treat osteoporosis.

Pang et al WO93/06845, published April 15, 1993, describes analogues of hPTH which involve substitutions of Arg²⁵, Lys²⁶, Lys²⁷ with numerous amino acids, including alanine, asparagine, aspartic acid, cysteine, glutamine, glutamic acid, glycine, histidine, isoleucine, leucine, methionine, phenylalanine, proline, serine, threonine, tryptophan, tyrosine or valine. These are claimed to be effective in the treatment of osteoporosis with minimal effects on blood pressure and smooth muscle.

The biological activity of hPTH is reflected in the activation of two second messenger systems, G-protein coupled adenylyl cyclase (cAMPase) and cAMPase coupled and uncoupled protein kinase C(PKC) activity. It has been established that the increase in bone growth, i.e. that effect which is useful in the treatment of osteoporosis, is coupled to the ability of the peptide sequence to increase cAMPase activity. The native hPTH containing only the first 34 amino acids has been shown to have all activities. It is typically shown as:

	Ser	Val	Ser	Glu	Ile	Gln	Leu	Met	His	Asn	
	1				5					10	
	Leu	Gly	Lys	His	Leu	Asn	Ser	Met	Glu	Arg	
					15					20	A
5	Val	Glu	Trp	Leu	Arg	Lys	Lys	Leu	Gln	Asp	
					25	26	27			30	
	Val	His	Asn	Phe							

It is the object of the present invention to produce new hPTH analogues having increased cAMPase activity with minimal side effects.

Brief Summary of the Invention

The above molecule "A" may have either a free carboxyl or amide ending in the sequence.

According to one feature of the present invention, it has been found that the cAMPase independent PKC is restricted to the 28-34 region of the above molecule "A". On the other hand, cAMPase activity has been shown to require the first 30 residues of the molecule. Thus, in accordance with this embodiment of the invention, it is possible to remove those biological activities associated with the cAMPase independent PKC activity by deleting a selected terminal portion of the hPTH-(1-34) molecule. In order for these shortened analogues to exhibit maximum activity, they must be in the form of the carboxyl terminal amides. One feature of the invention therefore comprises the human parathyroid analogues hPTH-(1-28-NH₂), hPTH-(1-29)-NH₂, hPTH-(1-30)-NH₂ and hPTH-(1-31)-NH₂.

According to another feature of the present invention, it has surprisingly been found that simply replacing Lys²⁷ with a Leu in the above molecule "A" is capable of increasing binding to the receptor and results in a higher activity for cAMPase stimulation. This analogue also exhibits its maximum activity only when in the form of the carboxyl terminal amide. Thus, another feature of the invention comprises human parathyroid analogues including all sequences from [Leu²⁷]-hPTH-(1-28)-NH₂ to [Leu²⁷]-hPTH-(1-34)-NH₂.

It is believed that this activity relating to Lys²⁷ is because of an amphiphilic α -helix near the carboxyl terminus of the above molecule is essential for the binding of hPTH to its receptor such as to stimulate cAMPase activity. This amphiphilic α -helix includes residues 20-34, with the most

stable helix being between residues 20 and 29. In receptor peptide complexes, where the bound peptide is in an α -helical conformation, the hydrophobic face is bound to the receptor. It is believed that Lys²⁷ is a single polar residue on the hydrophobic phase of this supposed helix and that the increased binding is achieved by replacing the Lys²⁷ with another amino acid.

A further feature of the invention comprises cyclic analogues based on the above molecule "A". These are in the form of lactams, along the polar face of the helix. As with the above features of the invention, the cyclic analogues are also most effective when in the form of carboxyl terminal amides and provide improved activity and/or stability. The cyclic analogues include cyclo (Lys²⁶-Asp³⁰) [Leu²⁷] -hPTH-hPTH-(1-34)-NH₂ and cyclo (Lys²⁷-Asp³⁰) -hPTH-(1-34)-NH₂.

Brief Description of the Drawings

Fig. 1 shows the structure of natural human PTH
(SEQ ID NO: 1)

Fig. 2 shows the structure of [Leu²⁷] -hPTH-(1-34)-NH₂
(SEQ ID NO: 2)

Fig. 3 shows the structure of hPTH-(1-31)-NH₂
(SEQ ID NO: 3)

Fig. 4 shows the structure of [Leu²⁷] -hPTH-(1-31)-NH₂
(SEQ ID NO: 4)

Fig. 5 shows the structure of hPTH-(1-30)-NH₂
(SEQ ID NO: 5)

Fig. 6 shows the structure of [Leu²⁷] -hPTH-(1-30)-NH₂
(SEQ ID NO: 6)

Fig. 7 shows the structure of cyclo (Lys²⁶-Asp³⁰) [Leu²⁷] -
hPTH-(1-34)-NH₂ (SEQ ID NO: 2)

Fig. 8 shows the structure of cyclo (Lys²⁷-Asp³⁰) -hPTH-
(1-34)-NH₂ (SEQ ID NO: 1)

Fig. 9 shows the structure of hPTH-(1-29)-NH₂

Fig. 10 shows the structure of hPTH-(1-28)-NH₂

Fig. 11 shows trabecular and cortical mass data for hPTH analogues of the invention, and

Fig. 12 shows further trabecular and cortical mass data

of the invention.

Description of the Preferred Embodiments

The structure of human parathyroid hormone (hPTH) is shown in Fig. 1 (SEQ ID NO: 1). Representative synthetic analogues are described in Table 1 and are further shown in Fig. 2-10 and SEQ ID NO: 2 - SEQ ID NO: 8. Trabecular and cortical mass data for various of these analogues is shown in Figs. 11 and 12.

Preparation of Hormone Analogues

The technique of solid phase synthesis developed by R.B. Merrifield ("Solid-Phase Peptide Synthesis", Advanced in Enzymology, 32, 221-296 (1969)), is widely and successfully used for the synthesis of polypeptides such as parathyroid hormone. The strategy is based on having the carboxyl-terminus amino acid of the support attached to a solid support. Successive amino acids are then added in very high yield. The α -amino group is protected in such a way that this protecting group can be removed without the removal of groups protecting side-chain reactive groups of the peptide from the solid support. The chemistry used here involves a modification of the original Merrifield method, referred to as the Fmoc approach. The Fmoc (fluorenyl-methoxycarbonyl) group can be removed by mild alkaline conditions which leave the alkali stable side-chain protecting groups and the link to the support untouched. This technique is described by E. Atherton & R.C. Sheppard (1989), "Solid Phase Peptide Synthesis: a Practical Approach", IRL Press, New York, N.Y.

Example 1

Synthesis & Purification of Linear hPTH Analogues

The α -amino groups of the amino acids were protected with 9-Fluorenyl-methoxycarbonyl (Fmoc) during coupling. Couplings were performed with a mixture of hydroxybenzotriazole (HOBT), 2-(1H-Benzotriazole-1-yl)-1, 1, 3, 3-tetramethyluronium tetrafluoroborate (TBTU), and diisopropylethylamine (DIPEA). A 4-fold excess of activated amino acids was used with double coupling on addition of the Asn, Gln, His, Val, and Ile

residues. The coupling times for Arg and Gly additions were increased from 30 min. to 60 min. Side chain protection of the amino acids was provided by: (1) the ϵ -amino group of lysine was protected as the 2-chlorobenzyloxycarbonyl derivative; (2) the guanido group of arginine was protected as the methoxytrimethylphenylsulfonyl derivative; (3) the carboxyl groups of glutamic and aspartic acids were protected as the t-butyl esters; (4) the hydroxyl group of serine was protected as the t-butyl ether; (5) the imidazole nitrogen of histidine was protected as the trityl derivative; (6) the amide nitrogens of glutamine and asparagine were protected as the trityl derivatives. Amino acid derivatives were purchased from Bachem Chemicals California. Analogues were synthesized with a Milligen 9050 Plus continuous-flow peptide synthesizer on TentaGel S-RAM as the solid support. Cleavage from the support yielded the terminal carboxylamide derivative as the product. Syntheses to yield the free carboxyl terminus were carried out with the appropriate Fmoc amino acyl derivative of Nova Syn TGA resin (Nova Biochemicals). For example, the synthesis of [Asp³⁵]-hPTH-(1-35)-COOH began with Fmoc-Asp(OtBu) Nova Syn TGA as the support.

Simultaneous cleavage from the solid support was carried out with 95% TFA in the presence of 6% thioanisole, 6% phenol, and 3% 1,2-ethanedithiol. The crude peptide was then precipitated with diethylether and lyophilized.

The crude product were purified by HPLC on a semi-prep PLRP-S column (Polymer Laboratories) (7.5 x 300 mm, 10 μ m), using 1%/min gradient of 0.1% trifluoroacetic acid in acetonitrile into 0.1% trifluoroacetic acid in water.

Example 2

Synthesis and Purification of Cyclic Analogues

Cyclo(Lys²⁶-Asp³⁰) [Leu²⁷]-hPTH-(1-34)-NH₂. The synthesis was performed similarly to Example 1 except in the region of the lactam. The side chain carboxyl group of Asp³⁰ was protected temporarily as the t-butyl ester. The side chain amino group of Lys was protected as the t-butyloxycarbonyl derivative. The coupling of Lys²⁶ was accomplished outside of

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the synthesizer. The resin was swollen in dichloromethane and the t-butyl group of Asp³⁰ and t-butyloxycarbonyl group of Lys²⁶ were removed by treatment with 30% trifluoroacetic acid in dichloromethane for 15 min. The resin was washed with
 5 dichloromethane, dimethylformamide, and dichloromethane in turn, neutralized with 20% diisopropylethylamine, then washed again with dichloromethane, dimethylformamide, and dichloromethane. The peptide-resin was then returned to the synthesizer and cyclization was accomplished by 2 cycles of 3
 10 hr each of (benzotriazolyl)-N-oxy-pyrrolidinium phosphonium hexafluorophosphate (PyBOP)/diisopropylethylamine/dimethylformamide (1:1:1) in 8-fold excess. Completeness of the reaction was monitored by a ninhydrin assay, as described in Kaiser et al (1970), Anal. Biochem. 34, 595-598.

15 Following lactam formation, any free residual amino groups were capped using acetic anhydride and 2-(1H-Benzotriazole-1-yl)-1, 1, 3, 3-tetramethyluronium tetrafluoroborate/diisopropylamine in dimethylformamide.

Cyclo-(Lys²⁷-Asp³⁰)-hPTH-(1-34)-NH₂- The synthesis was
 20 carried out in an analogous manner to that described above for cyclo (Lys²⁶-Asp³⁰) [Leu²⁷]-hPTH-(1-34)-NH₂, using OtBu protection of Asp-30 and t-butyloxycarbonyl protection of Lys²⁷

Example 3

Adenylyl Cyclase Assays

25 The ability of hPTH analogues to bind to receptors and activate the adenylyl cyclase coupled signalling mechanism was carried out on a differentiation competent osteoblast-like ROS 17/2 rat osteosarcoma cell line. Adenylyl cyclase activity was estimated by prelabelling the cellular ATP pool with [³H]-adenine and then measuring the amount of [³H]-cyclic AMP
 30 produced from the [³H]-ATP during the first 10 min. of exposure to a particular analogue. This was based on the procedures described by Whitfield et al (1992), J. Cell Physiol. 150, 299-303.

35 The adenylyl cyclase assay results are expressed in Table 1 below as the concentration necessary to express a half-maximal increase in the adenylyl cyclase activity.

Table 1

	Analogue	C50% max (adenylyl cyclase), nM
1	hPTH-(1-34)-NH ₂	15
2	[k271]-hPTH-(1-34)-NH ₂	6
3	HPH-(1-31)-NH ₂	20
4	[Leu ²⁷]-hPTH-(1-31)-NH ₂	14
5	HPH-(1-30)-NH ₂	21
6	[Leu ²⁷]-hPTH-(1-31)-NH ₂	15
7	c(Lys ²⁶ -Asp ³⁰)-[Leu ²⁷]-hPTH-(1-34)-NH ₂	7
8	c(Lys ²⁷ -Asp ³⁰)-hPTH-(1-34)-NH ₂	29
9	hPTH-(1-29)-NH ₂	28
10	hPTH-(1-29)-NH ₂	32

Example 5

5 Determination of Anabolic Activities of hPTH Analogues
with Ovariectomized Rat Model

A full description of the protocol is given in Rixon et al (1994), J. Bone, 9, 1179-1189. Sprague-Dawley rats weighing 255-275 g were purchased from Charles River (St. Constant, QC, Canada). For each experiment, 105 rats were weighed and divided into 21 groups, each with 5 rats, with comparable mean body weights between 260 and 265 g. These 21 groups were divided into 6 experimental groups consisting of 1 group of 5 animals for 0-time controls and 5 groups of 20 rats each which provided one group for normal or sham-ovariectomized (Sham-OVX) controls, one for OVX controls, and 3 for OVX rats treated with various hPTH analogues.

Sham OVX and OVX were performed under fluothane anaesthesia by the standard dorsal approach. For sham-OVX, the ovaries were exteriorized, but not removed. Except for the normal, unoperated rats, day 0 for each experimental group was the day of OVX. Starting 2 weeks later, designated groups of rats were given daily subcutaneously injections of PTH analogues (1 nmole/100g of body weight) dissolved in acidic saline (0.15 M NaCl containing 0.001 N HCl). The OVX control animals received comparable volumes of diluent solution only.

All animals survived the Sham- OVX and OVX operations, and there were no unscheduled deaths in any groups during the following 8 weeks.

5 The preparation and analysis of cortical and trabecular bone was carried out as described in M. Gunness-Hey & J.M. Hock (1984), Metab. Bone Dis. Rel. Res., 5, 177-181. Femurs were isolated, cleaned, and their lengths from the proximal, collum femoris to the distal condylar surfaces were measured. Each bone was then cut in half at mid-diaphysis and the
10 proximal half discarded. After removing the epiphysis, each half-femur was split lengthwise and the marrow washed out with distilled water. Each half was placed under a dissecting microscope and the trabecular (cancellous) bone was scraped out. The isolated trabecular bone and the remaining cortical
15 (compact bone) were dried at 55°C for at least 24 hr., and weighed to determined dry mass, expressed as mg/distal half-femur.

 After at least 3 days, the trichloroacetic acid extract was quantitatively removed and saved. The calcium contents of
20 the pooled trichloroacetic acid extracts from each cortical and trabecular bone sample were measured using the o-cresolphthalein complexone colorimetric procedure, using a kit from CIBA-Corning Diagnostics.

 The results obtained are shown in Figures 11 and 12.
25 Parts A and B of Figure 11 shows the trabecular (A) and cortical mass (B) data for various hPTH analogues tested in the ovariectomized rat model (Example 5), in which hPTH- (1-34) NH₂ (closed circles); sham OVX (open circles); hPTH-(1-31)-NH₂ (open triangles); OVX (open squares); hPTH-(8-84)-NH₂, a
30 PTH analogue lacking adenyl cyclase activity (closed squares).

 Further trabecular and cortical data is shown in parts A and B of Figure 12, in which hPTH-(1-34)-NH₂ (closed circles); sham OVX (open circles); OVX (open squares); hPTH-(1-30)-COOH,

5 nmols/100 g of rat dose (closed triangles); hPTH-(1-30)-COOH, 3 nmoles/100 g dose (open triangles).

The analogues of the present invention can be used in the treatment of osteoporosis and other bone related diseases and disorders involving bone cell calcium regulation.

The analogues of the present invention may be administered to a warm-blooded mammalian in need thereof, particularly a human, by parenteral, topical, rectal administration or by inhalation. The analogues may be conventionally formulated in a parenteral dosage form compounding about 1 to about 300 mg per unit of dosage with a conventional vehicle, excipient, binder, preservative, stabilizer, color, agent or the like as called for by accepted pharmaceutical practice.

For parenteral administration, a 1 to 10 ml intravenous, intramuscular or subcutaneous injection would be given one to four times daily. The injection would contain an analogue of the present invention in an aqueous isotonic sterile solution or suspension optionally with a preservative such as phenol or a solubilizing agent such as ethylenediaminetetraacetic acid (EDTA). Among the acceptable vehicles and solvents that may be employed are water, Ringer's solution and isotonic sodium chloride solution. In addition, sterile, fixed oils are conventionally employed as a solvent or suspending medium. Synthetic monoglycerides, diglycerides, fatty acids (such as oleic acid) find use as fixed oil in the preparation of injectables.

For rectal administration, the analogues of the present invention can be prepared in the form of suppositories by mixing with a suitable non-irritating excipient such as cocoa butter or polyethylene glycols.

For topical use, the analogues of the present invention can be prepared in the form of ointments, jellies, solutions, suspensions or dermal adhesive patches.

Daily doses are in the range of about 0.01 to about 200 mg per kg of body weight, depending on the activity of the specific compound, the age, weight, sex and conditions of the

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subject to be treated, the type and severity of the disease,
the frequency and route of administration. As would be well
known, the amount of active ingredient that may be combined
with the carried materials to produce a single dosage will
5 vary depending upon the host treated and the particular mode
of administration.

SEQUENCE LISTING

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- (ii) TITLE OF INVENTION: PARATHYROID HORMONE ANALOGUES
FOR THE TREATMENT OF OSTEOPOROSIS
- 10 (iii) NUMBER OF SEQUENCES: 8
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(F) POSTAL CODE: K1P 5P2
- (v) COMPUTER READABLE FORM:
20 (A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC Compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: Wordperfect 5.1
- (vi) CURRENT APPLICATION DATA:
25 (A) APPLICATION NUMBER:
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30 (A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
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- 35 (ix) TELECOMMUNICATION INFORMATION:
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- (2) INFORMATION FOR SEQ ID NO: 1:
- (i) SEQUENCE CHARACTERISTICS:
- 40 (A) LENGTH: 34 amino acids
(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

5	Ser	Val	Ser	Glu	Ile	Gln	Leu	Met	His	Asn	Leu	Gly
	1				5					10		
	Lys	His	Leu	Asn	Ser	Met	Glu	Arg	Val	Glu	Trp	Leu
			15					20				
10	Arg	Lys	Lys	Leu	Gln	Asp	Val	His	Asn	Phe		
	25					30						

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

	(A)	LENGTH:	34 amino acids
	(B)	TYPE:	amino acid
15	(D)	TOPOLOGY:	linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

20	Ser	Val	Ser	Glu	Ile	Gln	Leu	Met	His	Asn	Leu	Gly
	1				5					10		
	Lys	His	Leu	Asn	Ser	Met	Glu	Arg	Val	Glu	Trp	Leu
			15					20				
	Arg	Lys	Leu	Leu	Gln	Asp	Val	His	Asn	Phe		
	25					30						

25 (2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

	(A)	LENGTH:	31 amino acids
	(B)	TYPE:	amino acid
	(D)	TOPOLOGY:	linear

30 (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

	Ser	Val	Ser	Glu	Ile	Gln	Leu	Met	His	Asn	Leu	Gly
	1				5					10		
35	Lys	His	Leu	Asn	Ser	Met	Glu	Arg	Val	Glu	Trp	Leu
			15					20				
	Arg	Lys	Lys	Leu	Gln	Asp	Val					
	25					30						

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

5 (A) LENGTH: 31 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

10 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly
 1 5 10
 Lys His Leu Asn Ser Met Glu Arg Val Glu Trp Leu
 15 20
 Arg Lys Leu Leu Gln Asp Val
 25 30

15 (2) INFORMATION FOR SEQ ID NO: 5:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 30 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

25 Ser Val Ser Glu Ile Gln Leu Met His Asn Leu Gly
 1 5 10
 Lys His Leu Asn Ser Met Glu Arg Val Glu Trp Leu
 15 20
 Arg Lys Lys Leu Gln Asp
 25 30

(2) INFORMATION FOR SEQ ID NO: 6:

30 (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 30 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18. 19. 20. 21. 22. 23. 24. 25. 26. 27. 28. 29. 30. 31. 32. 33. 34. 35. 36. 37. 38. 39. 40. 41. 42. 43. 44. 45. 46. 47. 48. 49. 50. 51. 52. 53. 54. 55. 56. 57. 58. 59. 60. 61. 62. 63. 64. 65. 66. 67. 68. 69. 70. 71. 72. 73. 74. 75. 76. 77. 78. 79. 80. 81. 82. 83. 84. 85. 86. 87. 88. 89. 90. 91. 92. 93. 94. 95. 96. 97. 98. 99. 100. 101. 102. 103. 104. 105. 106. 107. 108. 109. 110. 111. 112. 113. 114. 115. 116. 117. 118. 119. 120. 121. 122. 123. 124. 125. 126. 127. 128. 129. 130. 131. 132. 133. 134. 135. 136. 137. 138. 139. 140. 141. 142. 143. 144. 145. 146. 147. 148. 149. 150. 151. 152. 153. 154. 155. 156. 157. 158. 159. 160. 161. 162. 163. 164. 165. 166. 167. 168. 169. 170. 171. 172. 173. 174. 175. 176. 177. 178. 179. 180. 181. 182. 183. 184. 185. 186. 187. 188. 189. 190. 191. 192. 193. 194. 195. 196. 197. 198. 199. 200. 201. 202. 203. 204. 205. 206. 207. 208. 209. 210. 211. 212. 213. 214. 215. 216. 217. 218. 219. 220. 221. 222. 223. 224. 225. 226. 227. 228. 229. 230. 231. 232. 233. 234. 235. 236. 237. 238. 239. 240. 241. 242. 243. 244. 245. 246. 247. 248. 249. 250. 251. 252. 253. 254. 255. 256. 257. 258. 259. 260. 261. 262. 263. 264. 265. 266. 267. 268. 269. 270. 271. 272. 273. 274. 275. 276. 277. 278. 279. 280. 281. 282. 283. 284. 285. 286. 287. 288. 289. 290. 291. 292. 293. 294. 295. 296. 297. 298. 299. 300. 301. 302. 303. 304. 305. 306. 307. 308. 309. 310. 311. 312. 313. 314. 315. 316. 317. 318. 319. 320. 321. 322. 323. 324. 325. 326. 327. 328. 329. 330. 331. 332. 333. 334. 335. 336. 337. 338. 339. 340. 341. 342. 343. 344. 345. 346. 347. 348. 349. 350. 351. 352. 353. 354. 355. 356. 357. 358. 359. 360. 361. 362. 363. 364. 365. 366. 367. 368. 369. 370. 371. 372. 373. 374. 375. 376. 377. 378. 379. 380. 381. 382. 383. 384. 385. 386. 387. 388. 389. 390. 391. 392. 393. 394. 395. 396. 397. 398. 399. 400. 401. 402. 403. 404. 405. 406. 407. 408. 409. 410. 411. 412. 413. 414. 415. 416. 417. 418. 419. 420. 421. 422. 423. 424. 425. 426. 427. 428. 429. 430. 431. 432. 433. 434. 435. 436. 437. 438. 439. 440. 441. 442. 443. 444. 445. 446. 447. 448. 449. 450. 451. 452. 453. 454. 455. 456. 457. 458. 459. 460. 461. 462. 463. 464. 465. 466. 467. 468. 469. 470. 471. 472. 473. 474. 475. 476. 477. 478. 479. 480. 481. 482. 483. 484. 485. 486. 487. 488. 489. 490. 491. 492. 493. 494. 495. 496. 497. 498. 499. 500. 501. 502. 503. 504. 505. 506. 507. 508. 509. 510. 511. 512. 513. 514. 515. 516. 517. 518. 519. 520. 521. 522. 523. 524. 525. 526. 527. 528. 529. 530. 531. 532. 533. 534. 535. 536. 537. 538. 539. 540. 541. 542. 543. 544. 545. 546. 547. 548. 549. 550. 551. 552. 553. 554. 555. 556. 557. 558. 559. 560. 561. 562. 563. 564. 565. 566. 567. 568. 569. 570. 571. 572. 573. 574. 575. 576. 577. 578. 579. 580. 581. 582. 583. 584. 585. 586. 587. 588. 589. 590. 591. 592. 593. 594. 595. 596. 597. 598. 599. 600. 601. 602. 603. 604. 605. 606. 607. 608. 609. 610. 611. 612. 613. 614. 615. 616. 617. 618. 619. 620. 621. 622. 623. 624. 625. 626. 627. 628. 629. 630. 631. 632. 633. 634. 635. 636. 637. 638. 639. 640. 641. 642. 643. 644. 645. 646. 647. 648. 649. 650. 651. 652. 653. 654. 655. 656. 657. 658. 659. 660. 661. 662. 663. 664. 665. 666. 667. 668. 669. 670. 671. 672. 673. 674. 675. 676. 677. 678. 679. 680. 681. 682. 683. 684. 685. 686. 687. 688. 689. 690. 691. 692. 693. 694. 695. 696. 697. 698. 699. 700. 701. 702. 703. 704. 705. 706. 707. 708. 709. 710. 711. 712. 713. 714. 715. 716. 717. 718. 719. 720. 721. 722. 723. 724. 725. 726. 727. 728. 729. 730. 731. 732. 733. 734. 735. 736. 737. 738. 739. 740. 741. 742. 743. 744. 745. 746. 747. 748. 749. 750. 751. 752. 753. 754. 755. 756. 757. 758. 759. 760. 761. 762. 763. 764. 765. 766. 767. 768. 769. 770. 771. 772. 773. 774. 775. 776. 777. 778. 779. 780. 781. 782. 783. 784. 785. 786. 787. 788. 789. 790. 791. 792. 793. 794. 795. 796. 797. 798. 799. 800. 801. 802. 803. 804. 805. 806. 807. 808. 809. 810. 811. 812. 813. 814. 815. 816. 817. 818. 819. 820. 821. 822. 823. 824. 825. 826. 827. 828. 829. 830. 831. 832. 833. 834. 835. 836. 837. 838. 839. 840. 84

| | | | | | | | | | | | | |
|---|-----|------|-------------------------------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | (xi) | SEQUENCE DESCRIPTION: SEQ ID NO: 6: | | | | | | | | | |
| | Ser | Val | Ser | Glu | Ile | Gln | Leu | Met | His | Asn | Leu | Gly |
| | 1 | | | | 5 | | | | | 10 | | |
| 5 | Lys | His | Leu | Asn | Ser | Met | Glu | Arg | Val | Glu | Trp | Leu |
| | | | 15 | | | | | 20 | | | | |
| | Arg | Lys | Leu | Leu | Gln | Asp | | | | | | |
| | 25 | | | | | 30 | | | | | | |

(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 29 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

| | | | | | | | | | | | | |
|----|-----------|-----|-----------|-----|----------|-----|-----|-----------|-----|-----------|-----|-----|
| 15 | Ser
1 | Val | Ser | Glu | Ile
5 | Gln | Leu | Met | His | Asn
10 | Leu | Gly |
| | Lys | His | Leu
15 | Asn | Ser | Met | Glu | Arg
20 | Val | Glu | Trp | Leu |
| 20 | Arg
25 | Lys | Lys | Leu | Gln | | | | | | | |

(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:

25 (A) LENGTH: 28 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

| | | | | | | | | | | | | |
|----|-----------|-----|-----------|-----|----------|-----|-----|-----------|-----|-----------|-----|-----|
| | Ser
1 | Val | Ser | Glu | Ile
5 | Gln | Leu | Met | His | Asn
10 | Leu | Gly |
| 30 | Lys | His | Leu
15 | Asn | Ser | Met | Glu | Arg
20 | Val | Glu | Trp | Leu |
| | Arg
25 | Lys | Lys | Leu | | | | | | | | |

What Is Claimed Is:

1. A human parathyroid hormone analogue (hPTH) selected from the group consisting of hPTH-(1-28)-NH₂, hPTH-(1-29)-NH₂, hPTH-(1-30)-NH₂ and hPTH-(1-31)-NH₂.
2. A human parathyroid hormone analogue selected from the group consisting of [Leu²⁷]-hPTH-(1-28)-NH₂, [Leu²⁷]-hPTH-(1-29)-NH₂, [Leu²⁷]-hPTH-(1-30)-NH₂, [Leu²⁷]-hPTH-(1-31)-NH₂, [Leu²⁷]-hPTH-(1-32)-NH₂, [Leu²⁷]-hPTH-(1-33)-NH₂, [Leu²⁷]-hPTH-(1-34)-NH₂.
3. A cyclic analogue of a human parathyroid hormone selected from the group consisting of cyclo(Lys²⁶-Asp³⁰) [Leu²⁷]-hPTH-(1-34)-NH₂ and cyclo (Lys²⁷-Asp³⁰)-hPTH-(1-34)-NH₂.
4. A human parathyroid hormone analogue (hPTH) or pharmaceutically acceptable salts thereof, selected from the group consisting of

hPTH-(1-28)-COOH
 hPTH-(1-29)-COOH
 hPTH-(1-30)-COOH
 hPTH-(1-31)-COOH
 [Leu²⁷]-hPTH-(1-28)-COOH
 [Leu²⁷]-hPTH-(1-29)-COOH
 [Leu²⁷]-hPTH-(1-30)-COOH
 [Leu²⁷]-hPTH-(1-31)-COOH
 [Leu²⁷]-hPTH-(1-32)-COOH
 [Leu²⁷]-hPTH-(1-33)-COOH
 [Leu²⁷]-hPTH-(1-34)-COOH
 cyclo (Lys²⁶-Asp³⁰) [Leu²⁷]-hPTH-(1-34)-COOH, and
 cyclo (Lys²⁶-Asp³⁰)-hPTH-(1-34)-COOH

5. Use of a human parathyroid hormone (hPTH) analogue according to Claim 1, 2, 3 or 4, for the treatment of osteoporosis in a warm-blooded animal in need of such treatment.
6. A composition for administration to a warm-blooded animal in need thereof, comprising a therapeutically effective amount of a human parathyroid hormone (hPTH) analogue according to Claim 1, 2, 3 or 4, in association with a pharmaceutically acceptable carrier or excipient.

Fig. 1 **2126299**

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Phe-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Ser-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-Gln-
Asp-Val-His-Asn-Phe-CONH₂

Fig. 2

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Leu-Leu-Gln-
Asp-Val-His-Asn-Phe-CONH₂

Fig. 3

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Phe-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-Gln-
Asp-Val-CONH₂

Fig. 4

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Phe-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Leu-Leu-Gln-
Asp-Val-CONH₂

Fig. 5

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-Gln-
Asp-CONH₂

Fig. 6

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-
Leu-Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Leu-Leu-Gln-
Asp-CONH₂

Fig. 7

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-Leu-
|
Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Leu-Leu-Glu-Asp-
|
Val-His-Asn-Phe-CONH₂

Fig. 8

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-Leu-
|
Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-Gln-Asp-
|
Val-His-Asn-Phe-CONH₂

Fig. 9

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-Leu-
Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-Gln-CONH₂

Fig. 10

H₂N-Ser-Val-Ser-Glu-Ile-Gln-Leu-Met-His-Asn-Leu-Gly-Lys-His-Leu-
Asn-Ser-Met-Glu-Arg-Val-Glu-Trp-Leu-Arg-Lys-Lys-Leu-CONH₂

2126299

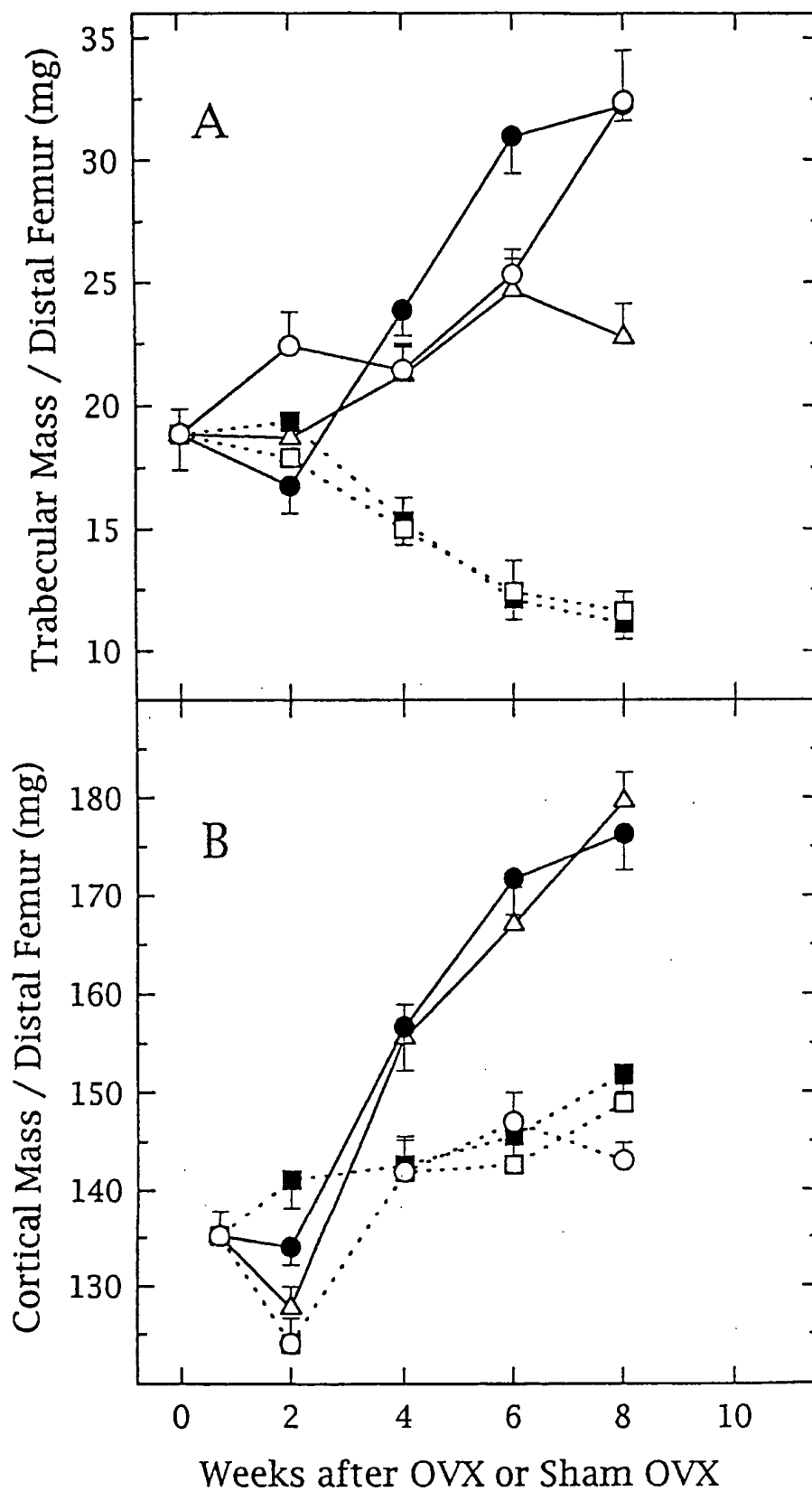
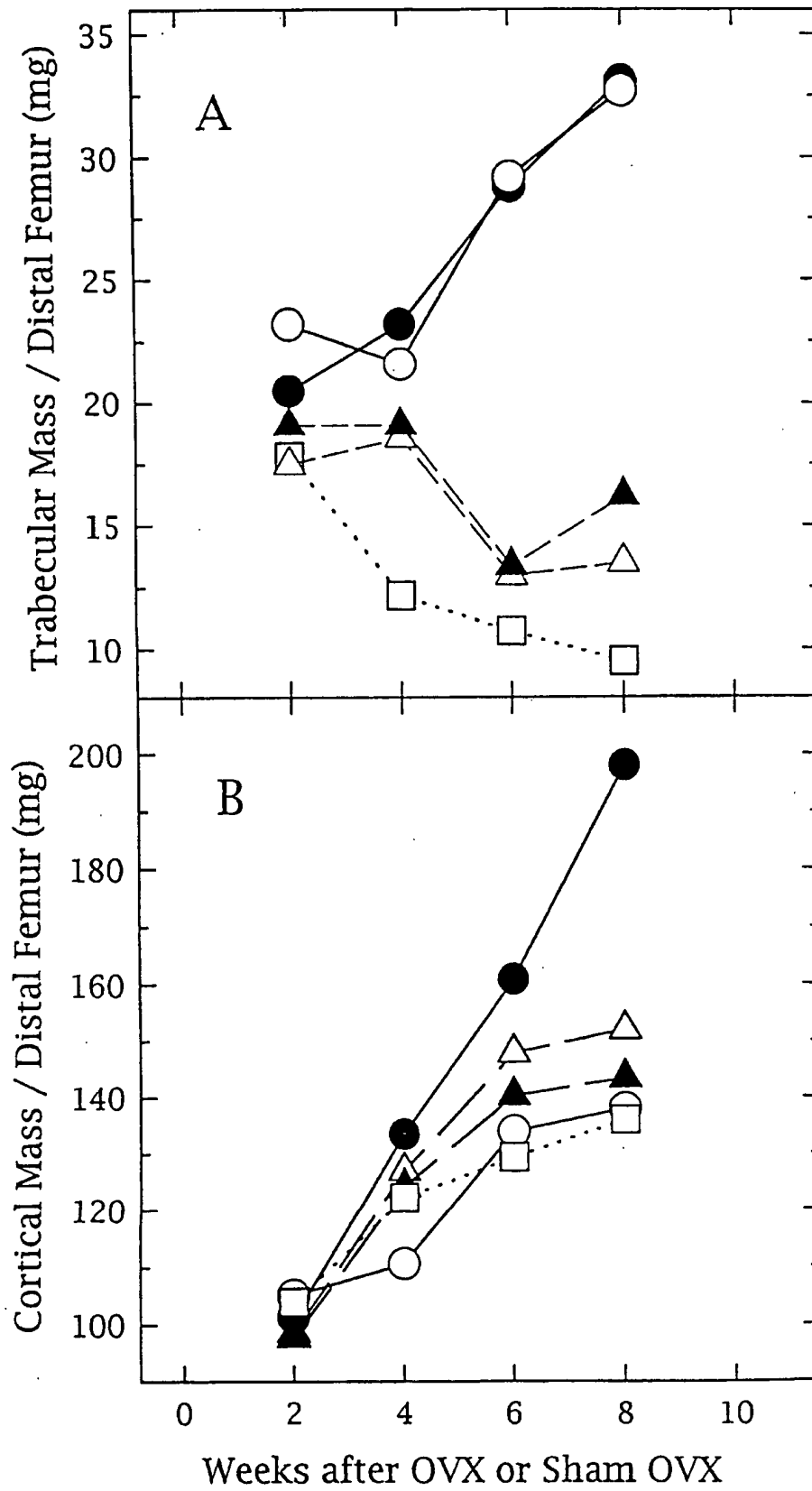


FIG. 11



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